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Uterine and placental distribution of selected extracellular matrix (ECM) components in the dog.

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Running title: ECM modification of canine pregnant uterus

Summary sentence: The canine uterine ECM is only moderately modified in early pregnancy, but undergoes vigorous reorganization processes in the uterus and placenta
20 following implantation.

Key words: dog (*Canis lupus familiaris*), extracellular matrix (ECM), collagen, connexins, TIMPs, pregnancy, remodeling, uterus.

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ABSTRACT

For many years, modifications of the uterine extracellular matrix (ECM) during gestation have not been considered as critical for successful canine (*Canis lupus familiaris*) pregnancy. However previous reports indicated an effect of free-floating blastocysts on the composition of the uterine ECM. Here, the expression of selected genes involved in structural functions, cell-to-cell communication and inhibition of matrix metalloproteinases were targeted utilizing qPCR and immunohistochemistry. We found that canine free-floating embryos affect gene expression of *FNI*, *ECM1* and *TIMP4*. This seems to be associated with modulation of trophoblast invasion, and proliferative and adhesive functions of the uterus. Although not modulated at the beginning of pregnancy, the decrease of structural ECM components (*i.e.*, *COL1*, -3, -4 and *LAMA*) from pre-implantation towards post-implantation at placentation sites appears to be associated with softening of the tissue in preparation for trophoblast invasion. The further decrease of these components at placentation sites at the time of parturition seems to be associated with preparation for the release of fetal membranes. Reflecting a high degree of communication, intercellular cell adhesion molecules are induced following placentation (*Cx26*), or increase gradually towards parturition (*Cx43*). The spatio-temporal expression of TIMPs suggests their active involvement in modulating of fetal invasiveness, and together with *ECM1* they appear to protect deeper endometrial structures from trophoblast invasion. With this, the dog appears to be an interesting model for investigating placental functions in other species, *e.g.*, in humans in which *Placenta accreta* appears to share several similarities with canine subinvolution of placental sites (SIPS).

INTRODUCTION

60 Interactions between the canine (*Canis lupus familiaris*) uterus and the corpora lutea (CL), and the interchange between the uterus and fetal trophoblast, represent remarkably unique mechanisms among mammalian species.

The bitch is a monoestric, non-seasonal breeder in which ovulation occurs spontaneously ([Concannon 1993](#)). As in other domestic animals CL are formed following ovulation, but in

65 contrast to other species, in canine pregnancy no placental steroid hormone synthesis arises ([Hoffmann et al. 1994](#)). Consequently, the increasing circulating steroid levels in both non-pregnant and pregnant bitches fully originate from the CL with luteal progesterone (P4) being responsible for establishment and maintenance of pregnancy until term ([Concannon et al. 1989](#)). Due to the lack of luteolysis in the absence of pregnancy, peripheral P4 levels are

70 almost identical in non-pregnant and pregnant bitches until shortly before parturition when prepartum luteolysis occurs signaling the onset of parturition ([Concannon et al. 1989](#)).

Consequently, in contrast to livestock, in the dog successful pregnancy does not depend on the inhibition of luteolysis to maintain the luteal life span and thereby embryo survival ([Kowalewski et al. 2009](#), [Gram et al. 2013](#), [Kowalewski 2014](#)). In other domestic animal

75 species pre-attachment embryos transmit signals needed to extend CL function ([Niswender et al. 2000](#), [Bazer et al. 2010](#), [Bazer 2015](#)), referred to as “maternal recognition of pregnancy” ([Short 1969](#)). This embryo-maternal communication synchronizes blastocyst development and uterine receptivity in a complex series of molecular and cellular mechanisms ([Seshagiri et al. 2009](#), [Paidas et al. 2010](#), [Geisert 2015](#)). In some species, e.g., in rodents or primates, the

80 uterine changes result in decidua formation and are therefore very pronounced. In primates, decidualization takes place even in the absence of embryos, while in rodents it is induced by their presence (reviewed in ([Geisert 2015](#))). In early pregnancy, the uterine extracellular matrix (ECM) influences trophoblast invasion ([Johnson et al. 2003](#)) and remodelling of the decidua ([Damsky et al. 1993](#), [Lala & Nandi 2016](#), [Smith et al. 2016](#)). The formation of

85 decidua, e.g., in mice, is associated with clearing of hyaluronic acid from the uterine stroma ([Brown & Papaioannou 1992](#)). In addition, *in vitro* studies demonstrate that decidualizing stromal cells of mouse and human origin produce ECM products such as basal lamina-like materials ([Wewer et al. 1985](#), [Wewer et al. 1986](#), [Kisalus et al. 1987](#)).

Despite the absence of an antiluteolytic signal at the beginning of pregnancy, in the dog

90 synchronization between blastocyst development and uterine preparation for implantation is also essential for healthy pregnancy. Thus, in dogs and other carnivores decidualization also takes place and is associated with implantation and placenta formation ([Wislocki & Dempsey](#)

1946, [Dempsey & Wislocki 1956](#), [Kautz et al. 2014](#), [Graubner et al. 2017b](#)). In accordance with the aforementioned *in vitro* decidualization studies with mouse and human cells, canine
 95 uterine cells can also be decidualized *in vitro*. Their response to a decidual stimulus (*i.e.*, increased secretory and proliferative activity) provides evidence that also in dogs uterine ECM modulation during decidualization occurs ([Kautz et al. 2015](#)). Furthermore, functional genomics studies imply that *in vivo* canine free-floating embryos transmit signals to the uterus and discretely, *i.e.*, without clearly visible morphological changes, affect uterine ECM
 100 composition at the early stage of pregnancy ([Graubner et al. 2017a](#)). Indicating the beginning of canine decidualization, these embryo-derived signals control the expression of uterine biochemical markers of decidualization ([Kautz et al. 2014](#)). The first morphological signs of decidualization are found later, commencing with attachment of the trophoblast ([Graubner et al. 2017b](#)). Ultimately, the uterine modifications become advanced and maternal-derived
 105 decidual cells are formed as an indispensable component of the endotheliochorial placenta in the dog. Decidual cells are needed for successful pregnancy since they are the only cells in the canine placenta expressing the nuclear P4 receptor (PGR) ([Vermeirsch et al. 2000](#), [Kowalewski et al. 2010](#)). Additionally, these cells and maternal endothelium can resist trophoblast digestion. Adequate embryo-maternal contacts between the trophoblast and
 110 maternal decidual cells, all embedded in the ECM components, are important for maintenance and termination of pregnancy. Thus, withdrawal of P4 from decidual cells will interfere with their physiological function and lead to abortion or preterm parturition ([Kowalewski et al. 2010](#)). ECM components appear to be involved here in feto-maternal communication leading to the luteolytic cascade, expulsion of fetuses and fetal membranes. Disturbances in embryo-
 115 maternal communication may result in clinical conditions like SIPS (subinvolution of placental sites), resembling placenta accreta in humans characterized by excessive trophoblast invasion.

Formation of the endotheliochorial placenta in the dog and other carnivores, indeed represents a unique type of placentation, with strong, but yet restricted, invasion. So far, little attention
 120 has been paid to uterine function in dogs during pregnancy, resulting in limited knowledge about its possible involvement in the establishment, maintenance and termination of pregnancy.

Because of this lack of information, driven by the hypothesis that ECM compounds are strongly modified throughout canine gestation, this study focused on the localization (utilizing
 125 immunohistochemistry (IHC)) and expression (using semi-quantitative TaqMan PCR) of several ECM proteins in the canine uterus. Selected stages of the reproductive cycle were

considered, starting with morpho-functional modifications induced by the presence of pre-attachment embryos, through implantation, early and late placentation, until prepartum luteolysis. A variety of different factors collectively represent the term ECM. Here, the
130 uterine localization and expression of ECM components with structural and adhesive functions were investigated (*e.g.*, collagen (COL) 1, -3 and -4, alpha smooth muscle actin (α SMA), and fibronectin (FN) 1). Additionally, proteins of the basal lamina, proteins involved in cell-to-cell communication, and proteins involved in inhibition of matrix metalloproteinases were targeted (*e.g.*, connexin (Cx) 26 and -43, laminin alpha (LAMA) 2,
135 extracellular matrix protein 1 (ECM1), and tissue inhibitor of metalloproteinases (TIMP) 2 and -4).

MATERIALS AND METHODS

Sample collection

140 For the present study, tissue material from 31 crossbred healthy bitches aged 2-8 years was used. Animals were randomly assigned to the following groups: (1) early pregnant (pre-attachment days 10-12; n=10); (2) corresponding non-pregnant controls (n=8); (3) early post-implantation (days 18-25 of pregnancy; n=5); (4) mid-gestation (days 35-40 of pregnancy; n=5); and (5) prepartum luteolysis (n=3).

145 Estrus detection was performed by cytological evaluation of vaginal smears and regular measurements of P4 concentrations every 2-3 days starting with the proestrous bleeding. The day when P4 blood levels increased > 5ng/ml for the first time was determined as the day of ovulation. Considering the peculiar need for canine oocyte maturation within the oviduct (on average 2-3 days after ovulation), bitches were mated 2 days after ovulation (defined as day 0
150 of gestation). Tissue collection was performed through routine ovariohysterectomy (OHE). In the early pregnant group (group 1), OHE was performed between days 10-12 of gestation. Pregnancy was confirmed by flushing free-floating (*i.e.*, pre-attachment) embryos. Dogs negative for embryo flushing were allotted to group 2 (corresponding non-pregnant control at day 10-12 of the luteal phase). The prepartum luteolysis takes place in the dog 12-24 h before
155 any visible physiological signs of parturition. Therefore, for group 5, blood samples were collected every 6 h starting on day 58 of pregnancy. When circulating P4 levels continued to decrease below 2-3 ng/ml in 3 consecutive measurements, indicating the prepartum luteolysis, samples were collected. Tissue collection in groups 1 and 2 consisted of uterine cross-sections (the whole thickness of the uterine wall). For groups 3-5, sampling included utero-placental
160 cross-sections (Ut-Pl; the whole thickness of the uterine wall, *i.e.*, uterus with adjacent

placenta). Additionally, for groups 3 and 4 inter-placental uterine cross-sections (inter-Pl; full thickness like groups 1 and 2) were collected. No inter-Pl samples were available from the natural luteolysis group of dogs (group 5).

For semi-quantitative real-time PCR, cellular RNA was stabilized by immersing the samples for 24h at 4°C in RNeasy Lysis Buffer (Ambion Biotechnologie GmbH, Wiesbaden, Germany); prolonged storage was at -80°C. For immunohistochemistry (IHC) studies, tissues were fixed for 24h in 10% neutral phosphate-buffered formalin at 4°C, and subsequently washed for 1 week with PBS (phosphate buffered saline) and embedded in paraffin wax. For additional information about tissue sampling and processing procedures, see previous reports (Kowalewski *et al.* 2010, Gram *et al.* 2013, Kautz *et al.* 2014). All uterine tissue samples were used for semi-quantitative real-time PCR, while 3 randomly chosen samples from each group were used for IHC.

All experimental procedures were carried out in accordance with animal welfare legislation and approved by the respective authorities of the University of Ankara (permits no. 2006/06 and 2008-25-124) Ankara, Turkey, and Justus-Liebig University Giessen (permit no. II 25.3-19c20-15c GI 18/14 and VIG3-19c-20/15c GI 18,14) Giessen, Germany.

RNA isolation, reverse transcription (RT), semi-quantitative (TaqMan) PCR and evaluation of data

Semi-quantitative TaqMan PCR was performed on total RNA isolated from all dogs. For this, TRIzol[®] reagent (Invitrogen, Carlsbad, CA, USA) was used according to the manufacturer's instructions. From each sample 10ng total RNA were used for RQ1 RNase-free DNase treatment (Promega, Dübendorf, CH). The RT reaction was performed according to the manufacturer's instructions with the High Capacity cDNA Reverse Transcription Kit including RNase Inhibitor (Applied Biosystems from Thermo Fisher Scientific, Foster City, CA, USA). Following this, amplification of cDNA was performed with the TaqMan[®] PreAmp Master Mix Kit (Applied Biosystems) according to the supplier's protocol. Detailed information about the TaqMan PCR procedure has been previously published (Kowalewski *et al.* 2006b, Kowalewski *et al.* 2011, Kautz *et al.* 2015). In short, reactions were run in duplicates with the Fast Start Universal Probe Master (ROX[®]) (Roche Diagnostics AG, Rotkreutz, CH) in an automated fluorometer (ABI PRISM[®] 7500 Sequence Detection System, Applied Biosystems). For negative controls, the so-called RT-minus control was used to check for potential genomic DNA contamination. Additional controls consisted of running experiments with autoclaved water instead of cDNA (no-template control) (Kowalewski *et al.*

195 [2006b](#), [Kowalewski et al. 2011](#)). If canine-specific TaqMan systems were not available commercially, Primer Express Software v 2.0 (Applied Biosystems) was used to select primers and probes which were purchased from Microsynth, Balgach, CH. TaqMan probes were designed with 6-FAM- and TAMRA-labels. For a list of self-designed primer-sequences and TaqMan systems see [Table 1](#). The self-designed expression assays were validated by the
200 CT slope method, which tests the TaqMan efficiency at different quantities of the targeted RNA as previously described ([Kowalewski et al. 2011](#)). The reaction efficiency for all self-designed assays was approximately 100%. Three reference genes, *GAPDH*, *Cyclophilin A* (*PPIA*) and *ACTIN-B*, were used for normalization to ensure homogeneity of variances. The following canine-specific TaqMan Gene Expression Assays were ordered from Applied
205 Biosystems: *PPIA* (Prod. No. Cf03986523-gH); *ACTIN-B* (Prod. No. Cf03023880_g1), collagen type 1, alpha 1 (*COL1A1*) (Prod. No. Cf02741575_mH); COL type 3, alpha 1 (*COL3A1*) (Prod. No. Cf02631366_m1); COL type 4, alpha 1 (*COL4A1*) (Prod. No. Cf02696157_mH). For relative quantification, the comparative CT method ($\Delta\Delta CT$ method) was applied as previously described ([Kowalewski et al. 2010](#), [Kowalewski et al. 2011](#)). The
210 sample with the lowest expression was used as a calibrator.

Statistical differences between the non-pregnant and early pregnant group (groups 1 and 2) were determined by an unpaired, two-tailed Student's t-test; a p-value <0.05 was considered as statistically significant. When target gene expression was compared in more than 2 observational groups (time-dependent expression at selected time points during pregnancy), a
215 parametric one-way ANOVA was applied. In the case of p<0.05, the Tukey-Kramer multiple comparisons post-test was performed. Statistical assumptions, such as normality and equality of variances were tested prior to applying the t-test or ANOVA. All analyses were performed with GraphPad 3.06 software (GraphPad Software, San Diego, CA, USA). Numerical data are presented as geometric means $X_g \pm$ geometric standard deviation (SD).

220 Immunohistochemistry

Standard immunohistochemistry (IHC) was performed to identify the localization of several ECM-proteins. Detailed information about our indirect immunoperoxidase method has been published previously ([Kowalewski et al. 2006a](#), [Kowalewski et al. 2006b](#)). In short, following
225 embedding in paraffin, tissues were sectioned into 2-3 μ m thickness, mounted onto SuperFrost microscope slides (Menzel-Glaser, Braunschweig, DE). Following de-paraffinization using xylene and rehydration in an ethanol series, antigen retrieval was performed using either heat-induced epitope retrieval (HIER) or proteolytic-induced epitope retrieval (PIER). HIER

was performed in a microwave oven at 560 W for 15 min in 10mM citrate buffer with pH 6.0.

230 PIER was performed using 0.25% pepsin (P7000-25G Sigma-Aldrich Chemie GmbH, Buchs, CH) in 10mM HCl at 37°C. The duration of PIER varied from 5 min to 90 min depending on the antibody (for detailed information see [Table 2](#)). Following antigen retrieval, sections were treated with 0.3% hydrogen peroxide in methanol for 30 min to block endogenous peroxidases. After that, blocking was performed with 10% normal serum from the same species in which the secondary antibody was produced, and overlaid with primary antibodies
235 overnight at 4°C. For negative controls, non-immune IgGs of the same species instead of the primary antibody at the same protein concentration as for the primary antibody were used (isotype controls).

Following incubation with the primary antibody, sections were washed with IHC buffer/0.3% Triton X pH 7.2-7.4 (0.8 mM Na₂HPO₄, 1.74 mM KH₂PO₄, 2.68 mM KCl, 137 mM NaCl),
240 and incubated with biotin-labeled secondary antibodies at 1:100 dilution. Next, the signals were enhanced using the Vectastain ABC Kit (Vector Laboratories, Inc., Burlingame, CA, USA) for 30 min, and immune reactions were visualized with the Liquid DAB+ substrate kit (Dako Schweiz AG, Baar, CH). Counter-staining was performed with hematoxylin. Next,
245 dehydration was performed in an ethanol series and xylene, and mounted using Histokit® (Assistant, Osterode, DE). Detailed information about commercially available primary and secondary antibodies, antigen retrieval and concentrations is given in [Table 2](#). All groups within one experiment were simultaneously stained with Liquid DAB+ substrate kit (Dako) for the same target using the same development time.

250 All slides were qualitatively evaluated for localization of the respective target protein and representative pictures were taken. Two researchers with good experience in canine uterine and placental histology evaluated the signals of target genes independently, using an ordinal system of none (-), weak (+/-), strong (+), and very strong (++) present ([Table 3](#)).

To identify the specific localization of TIMP2 und TIMP4 in the canine placenta, 4
255 consecutive slides (2-3µm thickness) were prepared from placenta in the prepartum luteolysis group. Each consecutive slide was stained with a different primary antibody (TIMP2, TIMP4, vimentin (VIM), pan-cytokeratin (pan-CYT)). Vimentin was used as a marker of mesenchymal cells in order to differentiate between different placental cellular compartments. It stains positively for decidual cells, endothelial cells and fibroblasts. Cytokeratin served as a
260 marker for epithelial cells, staining positively in fetal trophoblast. The staining and evaluation procedures were as described above.

RESULTS

mRNA expression of ECM-proteins in canine uterus, inter-Pl and Ut-Pl compartments at selected time points during pregnancy and prepartum luteolysis.

In all samples, the mRNA expression of ECM-proteins, cell adhesion molecules and modulators of ECM was detectable by applying semi-quantitative (TaqMan) PCR.

To evaluate embryo-induced changes in the uterus, a pairwise comparison between non-pregnant and early pregnant (pre-attachment) bitches was performed. In this contrast, *FNI* was significantly suppressed ($p=0.01$) by the presence of embryos during early pregnancy (pre-attachment), in contrast to increased *ECM1* and *TIMP4* expression ($p=0.01$) (Figure 1). *COL1*, *COL3*, *COL4*, *Cx26* and *-43* were not affected at the beginning of early pregnancy ($p>0.05$, not shown). Effects of the presence of embryos on *LAMA2* and *TIMP2* mRNA expression in the pre-implantation uterus were investigated previously (Graubner et al. 2017a) and were, therefore, excluded from the present study.

Next, the uterine pre-implantation stage was compared to inter-Pl sites and to the Ut-Pl compartments from different stages of pregnancy (i.e., post-implantation, mid-gestation and prepartum luteolysis). In inter-Pl sites uterine gene expression of *ECM1* was significantly upregulated at mid-gestation compared with pre- and post-implantation ($p<0.001$); it did not differ at earlier stages of pregnancy ($p>0.05$), i.e., pre- and post-implantation (Figure 2). The Ut-Pl compartments showed increased expression of the *ECM1* gene in the prepartum luteolysis group compared with pre-implantation uterus and Ut-Pl sites at early post-implantation stage ($p<0.001$ and $p<0.01$, respectively (Figure 2).

The uterine expression of all three major collagens, *COL1*, *COL3* and *COL4*, showed highest mRNA abundance at mid-gestation compared with either pre-implantation ($p<0.05$ for *COL1*, $p<0.01$ for *COL3* and $p<0.001$ for *COL4*) or post-implantation ($p<0.05$ for *COL1* and *COL4*). Whereas *COL1* and *COL4* did not differ between pre-implantation and post-implantation, *COL3* was higher in the early post-implantation group ($p<0.05$) (Figure 2). As for Ut-Pl compartments, early implantation was associated with suppression of *COL1* and *COL3* levels ($p<0.001$ for *COL1* and $p<0.05$ for *COL3*). Whereas they did not differ from post-implantation to mid-gestation, further suppression of both *COLs* mRNA expression was observed at prepartum luteolysis ($p<0.001$ and $p<0.01$ when compared with mid-gestation for *COL1* and *COL3*, respectively) (Figure 2). In contrast, the expression of *COL4* in the Ut-Pl comparison was unaffected at all examined stages of pregnancy ($p>0.05$).

FNI showed similar distribution patterns at inter-Pl sites as *COL1*, *COL3* and *COL4*. Following a more or less stable expression at inter-Pl at early gestation (i.e., during pre-

implantation and post-implantation), its uterine mRNA levels were highest at mid-gestation compared with previous gestational stages ($p<0.05$ and $p<0.001$ for pre-implantation and post-implantation, respectively) (Figure 3). At placentation sites (Ut-Pl compartments), the prepartum luteolysis was associated with downregulation of *FNI* compared with mid-gestation ($p<0.001$) (Figure 3).

The uterine mRNA levels of *LAMA2* were not significantly changed ($p>0.05$) during pre-implantation and post-implantation at the inter-Pl sites, but decreased thereafter, from post-implantation to mid-gestation ($p<0.001$) (Figure 3). *LAMA2* was strongly modulated at implantation sites, decreasing at Ut-Pl following early implantation ($p<0.001$), but it was not further changed towards prepartum luteolysis (Figure 3).

The uterine levels of *Cx26* were lowest at pre-implantation. They increased, however, significantly ($p<0.001$) at inter-Pl sites, showing the highest mRNA abundance post-implantation (Figure 3). Although apparently decreased at mid-gestation, *Cx26* levels were still more highly represented at mid-gestation compared to pre-implantation ($p<0.001$). When compared with Ut-Pl, the expression of *Cx26* was also significantly increased at implantation sites ($P<0.001$) and maintained its high mRNA levels until mid-gestation. The decrease towards prepartum luteolysis was significant compared with Ut-Pl during post-implantation and mid-gestation ($p<0.001$ and $p<0.001$, respectively) (Figure 3).

Uterine expression of the *Cx43* gene did not differ significantly at inter-Pl sites ($p>0.05$) (Figure 3). It was, however, significantly modulated in Ut-Pl compartments. Thus, implantation was associated with suppression of *Cx43* expression at placentation sites compared with the pre-implantation uterus ($p<0.05$). This expression continued, however, to increase at Ut-Pl gradually towards prepartum luteolysis, showing significant differences from post-implantation towards mid-gestation ($p<0.05$) and from mid-gestation to pre-partum luteolysis ($p<0.05$).

As for other genes, *i.e.*, *FNI*, *COL1*, -3, -4, the uterine mRNA encoding for *TIMP2* was highest at mid-gestation ($p<0.01$ and $p<0.001$ compared with pre-implantation and post-implantation, respectively) (Figure 4). *TIMP4* mRNA was already strongly induced at inter-Pl after implantation ($p<0.001$) and maintained its high levels through mid-gestation (Figure 4). At placentation sites, both *TIMP2* and *TIMP4* were strongly modulated and showed a gradual increase with progression of pregnancy, starting with early gestational stages until prepartum luteolysis when it was highest ($p<0.01$ and $p<0.05$ when compared with post-implantation) (Figure 4).

Localization of ECM proteins in the canine uterus and placenta

Localization and distribution patterns of COL1, COL3, COL4, FN1, ECM1, LAMA2, Cx26, Cx43, TIMP2 and TIMP4 were investigated at the protein level by applying IHC. All experimental groups were examined. A summary of the findings is presented descriptively in Table 3.

COL1 and COL3 were detectable in all maternal and fetal stromal compartments and in myometrium (Figure 5, Supplemental File 1, respectively). While at all compared stages of pregnancy (including non-pregnant controls at days 10-12), COL1 staining appeared to be evenly distributed, the intensity of COL3 was generally weaker compared with COL1.

COL4 was clearly detectable throughout gestation and was targeted to the basal lamina of blood vessels (Supplemental Files 2 D and G) in all compartments. Additionally, COL4 was detected in the myometrium of all groups with staining intensity varying, however, individually (Supplemental Files 2 B, F, I).

ECM1 was weakly detectable in luminal epithelium in the non-pregnant group (in one sample it was below the detection limit) (Figure 6 A and B). In early pregnancy, strong signals were detectable in luminal epithelium and in superficial and deep glands (Figures 6 C and D). During mid-gestation and at parturition luteolysis (Figure 6 E and H), it was only weakly detectable in the glandular epithelium of the superficial glands. At parturition luteolysis, ECM1 was abundantly present in the connective tissue layer separating superficial glands from deep glands (Figure 6 H). Following implantation (shown at mid-gestation) (Figure 6 F) and at parturition luteolysis (Figure 6 I), signals were detectable in glandular epithelium of deep glands. Placental localization of ECM1 was in cytotrophoblast (shown at mid-gestation and parturition luteolysis) (Figure 6 G and J).

Uterine endometrial expression of Cx43 was localized predominantly in epithelial and myometrial compartments (Figure 7). The luminal epithelium (Figure 7 A and C) and the epithelial cells of superficial and deep glands (Figure 7 A-F, H, I) stained strongly for Cx43. Additionally, Cx43 was present in the *tunica intima* of maternal blood vessels in all samples. In placenta (Figure 7 G and J), Cx43 was diffusely present in cyto- and syncytiotrophoblast and decidual cells. The strongest placental signals were found in the *tunica intima* of blood vessels (Figure 7 G and J).

Regarding TIMP2, it was only weakly expressed in the non-pregnant uterus (sometimes below detection limits) (Figure 8 A-B). In contrast, in the early pregnant uterus, at days 10-12 (prior to attachment), its expression was clearly detectable in the luminal epithelium, and superficial and deep glands (Figure 8 C, D). Myometrial signals were weak in both groups

([Figure 8 B, D](#)). Following implantation (shown at mid-gestation), endometrial TIMP2 was detectable in epithelial compartments of superficial and deep uterine glands ([Figure 8 E and F](#)); stromal compartments of the connective tissue layer appeared only weakly stained ([Figure 8 E](#)). At prepartum luteolysis, uterine TIMP2 signals were weaker in the epithelium of superficial glands ([Figure 8 H](#)). However, strong staining was observed in the connective tissue layer separating the superficial from deep glands ([Figure 8 H](#)). Myometrial signals were weak ([Figure 8 F, I](#)). Placental TIMP2 at mid-gestation was distributed ubiquitously and was found in cyto- and syncytiotrophoblast, in decidual cells and in vascular endothelial cells (*tunica intima*) ([Figure 8 G](#)). During prepartum luteolysis, placental expression was predominantly found in endothelial cells of maternal vessels ([Figure 8 J](#)). Sporadic nuclear staining was detected in the nuclei of myocytes, which did not allow further interpretation.

Compared with TIMP2, TIMP4 appeared to stain generally more weakly in all experimental groups and tissue compartments. Although weakly represented, uterine expression was localized in the luminal epithelium ([Figure 9 A and C](#)). Somewhat stronger staining was found in superficial and deep glands ([Figure 9 A-D, F](#)). Stronger TIMP4 signals were found during prepartum luteolysis in the connective tissue layer separating superficial from deep glands ([Figure 9 H](#)). It also appeared to be induced in deep uterine glands ([Figure 9 I](#)).

Within Ut-Pl compartments, placental TIMP4 was found in cytotrophoblast (shown at mid-gestation and during prepartum luteolysis; [Figure 9 G, J](#)). For Supplemental File 3, consecutive cross-sections of Ut-Pl compartment were prepared during prepartum luteolysis and were stained with either vimentin (VIM; [Supplemental File 3C](#)), pan-cytokeratin (pan-CYT; [Supplemental File 3D](#)), TIMP2 or TIMP4, allowing better differentiation between cellular tissue compartments stained by particular antibodies. In particular, localization patterns of both factors within canine placenta are clearly presented, with TIMP2 targeting to vascular endothelial cells and cytotrophoblast, presenting only weak signals in syncytiotrophoblast and maternal decidual cells ([Supplemental File 3A](#)). Clear TIMP4 signals were localized in cytotrophoblast ([Supplemental File 3B](#)).

In all samples, strong signals for aSMA were found in myometrium and the media of blood vessels ([Supplemental File 4](#)). Additionally, in the placenta aSMA was present in decidual cells, as shown at mid-gestation and prepartum luteolysis, ([Supplemental Files 4 G, J](#)).

FN1 was detectable throughout pregnancy. In general, stronger signals were found in epithelial than stromal compartments. The luminal epithelium of the non-pregnant group ([Supplemental File 5A](#)) was more strongly stained compared with the pre-attachment group ([Supplemental File 5C](#)). The myometrium stained weakly in all experimental groups and was

below detection limits in some samples ([Supplemental Files 5 B, F, I](#)). Additionally, diffuse
 400 signals for FN1 were observed throughout all cellular placental compartments, *i.e.*, syncytio-
 and cytotrophoblast, decidual cells and blood vessels ([Supplemental Files 5 G and J](#)).

LAMA2 ([Supplemental File 6](#)) was clearly present in all stromal compartments and in the
 media of blood vessels throughout pregnancy. Additionally, it was detectable in glandular
 epithelial cells in all groups. In the placental part of Ut-Pl compartments, LAMA2 staining
 405 was stronger in the stroma than in blood vessels ([Supplemental Files 6 G and J](#)).

Regarding Cx26, this protein was present in epithelial compartments throughout pregnancy
 ([Supplemental File 7](#)). Generally, signals were weaker in stromal and myometrial
 compartments. Cx26 was also detectable in endometrial epithelial cells. Following
 implantation, as shown during mid-gestation and at prepartum luteolysis, this protein was
 410 strongly locally induced in the epithelial cells of glandular chambers immediately above the
 connective tissue layer ([Supplemental Files 7 E, H](#)). Placental Cx26 was present diffusely in
 cyto- and syncytiotrophoblast and in decidual cells (indicated in [Supplemental File 6 J](#)).

DISCUSSION

415 The effects of modifications of uterine ECM on canine pregnancy have not been considered
 for many years. Because different tissues consist of dissimilar mixed combinations of cells
 which evolutionarily developed from different pedigrees, the configuration of ECM varies
 among tissues ([Abedin & King 2010](#)). These ECM components are involved in regulation of
 cell growth and differentiation, act as extracellular storage of hormones, and are involved in
 420 activation of signaling cascades (reviewed in: ([Hubmacher & Apte 2013](#), [da Anunciacao et al.
 2017](#))). In the uterus, the major ECM components are collagens, proteoglycans, hyaluronan
 and glycoproteins ([Leoni et al. 1990](#), [Oliveira et al. 2015](#), [Franczyk et al. 2017](#)). It is worth
 noting that in mammals the uterine ECM composition changes dynamically depending on the
 reproductive status (*i.e.*, non-pregnant vs. pregnant) ([Cabrol et al. 1985](#), [Leoni et al. 1990](#),
 425 [Hjelm et al. 2002](#), [Boos et al. 2003](#)), and in some species, such as humans and cattle, the
 composition of the uterine ECM is cycle-dependent ([Boos 2000](#), [Curry & Osteen 2001](#)). In
 fact, most details about uterine ECM modifications and functions during pregnancy are
 known only for humans and rodents. In these species, in early pregnancy the uterine ECM
 influences trophoblast invasion ([Johnson et al. 2003](#)) and remodeling of decidua ([Damsky et
 430 al. 1993](#), [Lala & Nandi 2016](#), [Smith et al. 2016](#)), while the ECM is modulated by matrix
 metalloproteinases (MMPs) of fetal (trophoblast) origin. As for the canine non-pregnant and
 pregnant uterus and placenta, the time- and organ-specific expression of MMP2 and -9 have

been investigated ([Beceriklisoy et al. 2007](#), [Fellows et al. 2012](#), [Diessler et al. 2017](#)). Their expression patterns indicate pregnancy-associated modifications of ECM.

435 The present study explored the spatio-temporal expression of ECM components selected from our preceding microarray analysis ([Graubner et al. 2017a](#)), and known for their involvement in modulating uterine and placental functions in other animals, in the canine uterus and placenta throughout pregnancy. Quantitative assessment was done by qPCR, and cellular localization was investigated qualitatively by IHC. The antibodies applied herein proved
440 unsuitable for western blot analysis. It appears plausible that linearization of proteins by reducing agents during western blot preparation affects the structure of target epitopes, thereby preventing antibody binding.

Structural and adhesive proteins of the ECM

445 Expression of the following proteins was assessed: COL1, COL3, COL4, ECM1, FN1, LAMA2 and aSMA. Free-floating embryos were previously reported to increase the uterine expression of *LAMA2* ([Graubner et al. 2017a](#)). Here, the uterine adjustments during early decidualization were additionally characterized by increased expression of *ECM1*, whereas *FN1* was suppressed. None of the major COLs was affected by the presence of embryos.
450 Showing distinctly different localization patterns and both being affected by presence of embryos, ECM1 and LAMA2 appear to be involved in the establishment of canine pregnancy. As a component of the lamina basalis, increased expression of LAMA is a marker of ongoing stromal decidualization in humans ([Church et al. 1997](#)). Similarly in the dog, even if devoid of apparent morphological differentiation of the uterus, pre-implantation embryos stimulated
455 the expression of LAMA2, predominantly in the uterine stroma. The increased ECM1 expression was abundantly localized in endometrial epithelium. In the human endometrium its expression during implantation is further stimulated by some of the chemokines critically involved in leukocyte migration and trophoblast invasion, *e.g.*, CX3CL1 and CCL14 ([Hannan & Salamonsen 2008](#)). Although not yet investigated, similar functional relationships cannot
460 be ruled out for the canine species. Further effects of ECM1 in the canine uterus might be associated with its proliferative activities. Thus, by interacting with the epidermal growth factor (EGF) related pathways, ECM1 promotes cellular differentiation and proliferation ([Lee et al. 2014](#)). Within the canine uterus, the localization patterns of EGF receptor (EGFR) were described previously ([Sagsoz et al. 2014](#)) and reflect the localization of ECM1 presented
465 herein. A possible functional relationship between these two entities is certainly worth attention and should be addressed in the future.

Regarding the embryo-induced suppression of the cell adhesion molecule *FN1*, similar to the dog, its decrease during early pregnancy also occurs in rodents ([Zollinger & Smith 2017](#)), a phenomenon that has been linked to the early decidualization process around the time of implantation ([Grinnell et al. 1982](#)). Moreover, reduction of FN1 expression seems to be important to prevent excessive trophoblast invasion into maternal tissues ([Kaloglu & Onarlioglu 2010](#)). Despite the less invasive type of canine placentation compared to rodents, the *FN1* decrease observed before trophoblast attachment in dogs could also possibly be linked to similar protective functions in order to control trophoblast invasion. A key mechanism in this context seems to be the cooperation of FN1 and integrin molecules in modulating trophoblast proteolytic activities [through/via] the ECM ([Sutherland et al. 1993](#), [Fazleabas et al. 1997](#)). Interestingly, however, the canine uterine expression of several integrins (ITG) (*e.g.*, ITGA2B, ITGB2 and ITGB3) increases in early pregnancy ([Bukowska et al. 2011](#)), while FN1 expression is diminished by the presence of embryos.

It needs to be emphasized that, in agreement with our previous findings ([Graubner et al. 2017a](#)), and in contrast to other species including humans and rodents, the above described functional changes in the canine uterus driven by the presence of embryos, take place without any morphologically visible alterations of uterine structures. The deep structural remodeling processes of the canine uterus start following implantation and placentation and are associated with increased expression of COLs (*COL1*, -3, and -4), *ECM1* and *FN1* in the uterine wall (*i.e.*, inter-PI). The increased expression of structural and basement membrane components, widely localized in stroma and, depending on the type of COL (*e.g.*, COL1 and -3), in myometrium, seems to relate to the uterine growth as a natural phenomenon resulting from the development of conceptuses.

Conversely, at implantation sites, trophoblast invasion and placentation decreased the expression of two major COLs (COL1 and -3) and LAMA2. This is apparently associated with the remodelling processes and placental development, as all COLs, and in particular the most robust COL1, were clearly detectable in utero-placental stromal compartments. Similar effects were observed in other animals displaying invasive types of placentation, *e.g.*, in rodents, in which implantation was associated with decreased *COL1* expression at invasion sites, implying the need for *COL1* suppression during remodeling events associated with embryo-induced decidualization and subsequent placentation in these animals ([Clark et al. 1993](#)).

The decrease in *LAMA2* found in our study additionally indicates loss of stability of the basal lamina, and associated with it migration and differentiation of cells ([Kleinman et al. 1985](#),

[Engvall et al. 1990](#), [Aumailley 2013](#)). This differentiates the dog from mammals with more invasive forms of placentation such as rodents and primates. Accordingly, around the time of implantation in the murine uterus, laminin is synthesized by decidual cells ([Wewer et al. 1985](#)) where it influences trophoblast migration ([Zhang et al. 2000](#), [Korgun et al. 2007](#)). In humans also decidual cells produce laminin ([Kisalus et al. 1987](#), [Aplin et al. 1988](#), [Haouzi et al. 2011](#)). At later stages of human gestation, several isoforms of laminin and fibronectin are found in the basal lamina of capillaries and in the stroma and trophoblast cells of the villi ([Korhonen & Virtanen 2001](#)). As for the dog, LAMA2 seemed to be more strongly represented in the stromal than epithelial compartments, in particular following implantation.

Besides laminin, the decidualization process in humans is marked by increased expression of aSMA ([Geisert 2015](#)). This protein seems to be strongly required during pregnancy since there is a relationship between abnormal actin polymerization and pregnancy failure in humans ([Montazeri et al. 2015](#)). In dogs, aSMA is only weakly present in uterine stromal cells of non-pregnant animals and at the pre-implantation stage. However, following placentation aSMA is strongly expressed in maternal decidual cells and can be used as a cellular marker of decidual cells in the canine species. As a matter of fact, aSMA was also strongly expressed in canine decidualized uterine primary stromal cells *in vitro* ([Kautz et al. 2015](#)).

Following attachment and invasion of the trophoblast in dogs the girdle placenta is developing. Histologically, at the placentation site a connective tissue layer can be found that separates enlarged superficial glands (so called glandular chambers) from deeper localized parts of the uterus. It functions as a barrier, protecting deep uterine glands and myometrium from proteolytic activity of the trophoblast. When this barrier is breached, an exaggerated invasion of trophoblast may result in SIPS ([Al-Bassam et al. 1981](#)). Clearly detectable staining of structural collagens (COL1, and -3) and basal lamina components such as LAMA2 was identified throughout pregnancy in the stroma of this connective tissue layer. Their respective mRNA levels were not modulated from post-implantation towards mid-gestation, but the major collagens, COL1 and -3, were strongly suppressed at the time of parturition luteolysis. This may be seen as an indicator of preparation for parturition and release of the fetal membranes (placentolysis). A similar role of physiological degradation of collagens during preparation for the release of fetal membranes has been discussed for cattle ([Attupuram et al. 2016](#)). In fact, a retained placenta can be successfully treated by administration of collagenases into the umbilical artery ([Eiler & Hopkins 1993](#), [Attupuram et al. 2016](#)).

535 With regards to *FNI*, which was generally more strongly represented in epithelial compartments, its uterine and placental expression followed the expression patterns exhibited by COLs. This leads to the assumption that the main role of *FNI* in the canine uterus is a bridging function between collagens and other ECM components.

The utero-placental *ECM1* expression increased with the progression of pregnancy and was
 540 highest at parturition luteolysis. Within the placenta, ECM1 stained predominantly in cytotrophoblast, which appeared to be the major source of ECM1 in the Ut-Pl compartment. This resembles results observed in humans where the importance of ECM1 during maintenance of pregnancy and fetal development is underlined by its highest expression in the placenta and fetal heart tissue but not in other organs (*e.g.*, brain, lung, liver, kidney, pancreas
 545 or skeletal muscles) ([Smits et al. 1997](#)). In addition to the aforementioned function of ECM1 in promoting proliferation and differentiation of cells during the onset of pregnancy, at prepartum luteolysis the co-localization of ECM1 with TIMP2- and -4 in the tissue layer separating superficial from deep uterine glands strongly attracted our attention. This spatio-temporal detection is specific to prepartum luteolysis, but not at earlier stages of pregnancy. A
 550 possible implication for its function arises from *in vitro* studies showing ECM1-mediated reduction of MMP9 proteolytic activity in a human model ([Fujimoto et al. 2006](#)). Cumulatively, a protective role of ECM1, possibly interacting with a metalloproteinases activity-balancing system, should be considered.

555 **Connexins (Cx26 and -43)**

Connexins (Cx) build gap junction proteins, through which two cells can exchange small molecules, electrical charges and second messengers ([Bruzzone et al. 1996](#), [Kumar & Gilula 1996](#), [Evans & Martin 2002](#)). The uterine expression of Cx26 and -43 has been described during the menstrual cycle in humans as well as during pregnancy in humans, rats and sheep
 560 ([Winterhager et al. 1993](#), [Grummer et al. 1994](#), [Jahn et al. 1995](#), [Johnson et al. 2017](#)). Both hormonal- and embryo implantation-mediated effects were observed ([Grummer et al. 2004](#)). Successful implantation of embryos in rodents requires a complete suppression of both connexins prior to implantation ([Grummer et al. 1994](#), [Grummer et al. 2004](#)). At the time of implantation connexins are re-induced ([Grummer et al. 1996](#)). The importance of Cx during
 565 establishment of pregnancy has been further strengthened by their abnormal expression patterns reported in pathologies such as recurrent pregnancy loss ([Laird 2006](#), [Nair et al. 2011](#)).

No such information was so far available regarding the canine uterus and placenta, which prompted us to investigate the expression and distribution patterns of Cx26 and -43 throughout pregnancy. The uterine expression of both Cx remained unaffected by the presence of free-floating embryos. The respective proteins were detectable, however, with Cx43 appearing more abundantly expressed than Cx26. Whereas being clearly detectable, Cx43 was more or less constantly present in the uterine wall; its increasing expression in the Ut-Pl compartment followed placental development and reached its highest levels during prepartum luteolysis. It is noteworthy that, although rather ubiquitously expressed, within the placenta Cx43 mostly targeted to the maternal endothelium. This expression pattern differed strongly from that observed for Cx26, which appeared to be in agreement with observations made in other species exhibiting invasive types of placentation. Thus, the uterine and placental expression of Cx26 was strongly induced following implantation. Its expression was predominantly localized in the endometrial tissues, both at the placentation sites and at inter-Pl. The prepartum luteolysis was associated with a strong decrease of the respective mRNA levels.

Regarding the possible functions of connexins within the placenta, in humans Cx43 appears indispensable for proper decidualization and uterine angiogenesis ([Laws et al. 2008](#)). Additionally, it has been shown *in vitro* that Cx43 is involved in regulating differentiation of cytotrophoblast to syncytiotrophoblast and plays roles in feto-maternal exchange ([Cronier et al. 2002](#)). Based on the localization pattern of Cx43 described above, in the dog its involvement in uterine and placental angiogenesis also seems likely, since the strongest signals were noticed in the intima and media of placental blood vessels. The involvement of Cx43 during canine decidualization cannot, however, be ruled out and merits further investigations.

Spatio-temporal expression of TIMP2 and TIMP4

When discussing the development and function of the canine endotheliochorial placenta, the mechanisms controlling the invasive behavior of trophoblast need to be considered. In general, the invasive properties of trophoblast are at least in part provided by MMPs (reviewed in ([Goldman-Wohl & Yagel 2002](#), [Cohen et al. 2006](#))). At parturition, MMPs are additionally associated with placental detachment, as shown in humans and cattle ([Eiler & Hopkins 1992](#), [Strauss 2013](#), [Menon et al. 2016](#)). In dogs, MMPs are present in trophoblast cells ([Beceriklisoy et al. 2007](#), [Fellows et al. 2012](#), [Diessler et al. 2017](#)). Biologically active inhibitors, *i.e.*, TIMPs, by interacting with MMPs promote cell growth, inhibit angiogenesis

and reveal both anti- and pro-inflammatory effects (reviewed in ([Stetler-Stevenson 2008](#), [Brew & Nagase 2010](#))). Although TIMP2 and TIMP4 are 50% identical in sequence, they still bear differences in specificity to MMPs ([Brew & Nagase 2010](#)). *In vitro* studies with human

605 endometrial cells have shown that TIMPs are positively associated with the decidualization process and cell migration ([Graham et al. 2017](#)). Interestingly, *TIMP2* expression can be induced by P4 ([Imada et al. 1994](#), [Jo et al. 2015](#)). We have previously shown the ability of canine embryos to increase uterine *TIMP2* expression prior to implantation ([Graubner et al. 2017a](#)). During canine labor also, the placental presence of *TIMP2* has been confirmed at the

610 mRNA level ([Fellows et al. 2012](#)). It is assumed that interference with TIMPs functions can lead to placental retention and SIPS in dogs, or to spontaneous early pregnancy failure and *Placenta accreta* in humans ([Al-Bassam et al. 1981](#), [Ke et al. 2006](#), [Nissi et al. 2013](#)). In addition to the previously reported increase in mRNA expression of *TIMP2* ([Graubner et al. 2017a](#)), here also, elevated TIMP4 expression was observed in the pre-implantation canine

615 uterus. The stimulatory effect on gene expression was supported by IHC results in which free-floating embryos appeared to increase signals for TIMP2 in both superficial and deep uterine glands. Following implantation, a gradual increase was noted for both TIMPs (*i.e.*, *TIMP2* and -4) in uterine and placental compartments.

An important finding from our study is the strong presence of TIMP2 and -4 in the

620 endometrial connective tissue layer separating the superficial from deep glands. In a healthy pregnancy, trophoblast invasion is stopped at this bordering protective structure. Additionally, uterine and placental TIMPs were localized around blood vessels. Thus, it is plausible to assume that both proteins are involved in the protection of maternal tissue from trophoblast invasion. Within the placenta, TIMP2 appeared more abundantly represented than TIMP4.

625 Both factors were, however, localized in trophoblast cells, which appears to reflect an auto-/paracrine feedback loop acting within these cells, possibly controlling own proteolytic function of trophoblast and protecting the maternal tissues from excessive invasion. Although it is not known for the dog if TIMP2 and TIMP4 interact with active MMP2 and MMP9, our findings (*i.e.*, the gradual increase of *TIMP2*, *TIMP4* and *ECM1*) indicate that MMP-

630 mediated-trophoblast invasion might be balanced by these factors. In this regard, future studies should be considered, investigating for example the expression of MMPs and their regulators within excessive trophoblast invasion leading to conditions such as SIPS.

Conclusions

635 In this study we investigated the modulation of uterine and placental ECM in defined stages
of canine pregnancy. Based on the presented results, it seems that the primary goal of the
early embryo maternal communication is the regulation of trophoblast invasion, and the
proliferative and adhesive functions of the uterus. Following trophoblast attachment, the ECM
is strongly modulated reflecting dynamic feto-maternal interactions during establishment of
640 the canine endotheliochorial placenta.

The role of the endometrial connective tissue layer as an active, and not only physical but also
a biochemical, barrier protecting maternal tissues from unrestrained trophoblast invasion, is
strongly implied. This is supported by the abundant local expression of ECM components
potentially actively involved in modulating the invasiveness of fetal cells.

645 Regulatory mechanisms involved in modulation of ECM components during establishment
and maintenance of gestation modes characteristic of invasive placentation types, appear to be
similar to what has been described for other mammalian species exhibiting similar, or even
more intense invasion by trophoblast (haemochorial placentation type). In particular, TIMPs
and ECM1 appear to be plausible candidates involved in regulating the establishment and
650 termination of pregnancy in the dog. With this, the dog could provide an interesting model for
investigating placental functions in other species, *e.g.*, in humans in which *Placenta accreta*
appears to share several similarities with canine SIPS.

Abbreviations:

655 aSMA: alpha-smooth muscle actin
COL: collagen
CL: corpus luteum
Cx: connexin
ECM: extracellular matrix
660 ECM1: extracellular matrix protein 1
FN1: fibronectin1
GAPDH: glyceraldehyde 3-phosphate dehydrogenase
HIER: heat initiated epitope retrieval
IHC: immunohistochemistry
665 inter-Pl: inter-placenta
LAMA2: laminin alpha 2
min: minutes
MMP: matrix metalloproteinase
OHE: ovariectomy

670 P4: progesterone
pan-CYT: pan-cytokeratin
PGR: nuclear progesterone receptor
PIER: protease initiated epitope retrieval
PPIA: *Cyclophilin A*
675 qPCR: quantitative real-time polymerase chain reaction
SD: standard deviation
SIPS: subinvolution of placental sites
TIMP: tissue inhibitor of matrix metalloproteinase
Ut-Pl compartment: utero-placental compartment
680 VIM: vimentin

Declaration of interest:

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported. All authors read and approved the final version of the manuscript.

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Author contribution statement:

FRG: involved in developing the concept of the study, experimental design, generating data, analysis and interpretation of data and writing of the manuscript. AB: knowledge transfer, critical discussion of data, editing of the manuscript. SA and IK: knowledge transfer, critical discussion of data, collection of tissue material. MPK: designed and supervised the project, involved in interpretation of the data, drafting and revising the manuscript. All authors read and approved the final manuscript.

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Figures, Tables and Supplemental Files:

(submitted as separated file)

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Figures, Tables and Supplemental Files:

Tables:

Table 1:

List of primers used for real-time (TaqMan) PCR.

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Table 2:

List of primary and secondary antibodies used for indirect immunohistochemistry (IHC).
PIER = protease initiated epitope retrieval; HIER heat initiated epitope retrieval

10 **Table 3:**

Tabular, descriptive presentation of localization patterns of extracellular matrix proteins (ECM) as determined by immunohistochemistry (IHC) in canine uterine and placental compartments. (-) = not present; (+/-) = weakly present; (+) = strongly present; (++) = very strongly present. COL = collagen, FN1 = fibronectin 1, ECM1 = extracellular matrix protein 1, LAMA2 = laminin alpha 2, Cx = connexin, TIMP = tissue inhibitor of metalloproteinase, aSMA = alpha-smooth muscle actin.

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Figures:

Figure 1:

20 Expression of selected extracellular matrix (ECM) genes in the canine pre-implantation uterus (days 10-12 of pregnancy) affected by the presence of free-floating embryos, compared with gene expression in their non-pregnant counterparts as determined by Real Time (TaqMan) PCR. *FN1*= fibronectin 1; *ECM1*= extracellular matrix protein 1, *TIMP4* = tissue inhibitor of metalloproteinases 4. An unpaired, two-tailed Student's t-test was applied. P<0.05 was defined as significant. Bars with different asterisks differ at P=0.01. Numerical data are presented as geometric means $\bar{X}_g \pm$ geometric standard deviation (SD).

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Figure 2:

Expression of extracellular matrix protein 1 (*ECM1*), collagen (*COL*) 1, -3 and -4 as determined by Real Time (TaqMan) PCR in the canine uterus during different times of pregnancy. Two comparisons are presented for each gene: First, the pre-implantation (Pre-imp.) stage was compared to the inter-placental sites during post-implantation (Post-imp.) and mid-gestation (Mid-gest.), and next to the utero-placental compartments (placentation sites) of post-implantation (Post-imp.), mid-gestation (Mid-gest.) and prepartum luteolysis. Asterisks

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35 indicate ★= $p<0.05$, ★★= $p<0.01$, ★★★= $p<0.001$. Numerical data are presented as geometric means $Xg \pm$ geometric standard deviation (SD).

Figure 3:

40 Expression of fibronectin 1 (*FNI*), laminin alpha 2 (*LAMA2*), connexin (*Cx*) 26 and -43 as determined by Real Time (TaqMan) PCR in canine uterus during different times of pregnancy. Two comparisons are presented for each gene: First, the pre-implantation (Pre-imp.) stage was compared to inter-placental sites during post-implantation (Post-imp.) and mid-gestation (Mid-gest.), and next to the utero-placental compartments (placentation sites) of post-implantation (Post-imp.), mid-gestation (Mid-gest.) and prepartum luteolysis. Asterisks
45 differ at ★= $p<0.05$, ★★= $p<0.01$, ★★★= $p<0.001$. Numerical data are presented as geometric means $Xg \pm$ geometric standard deviation (SD).

Figure 4:

50 Expression of tissue inhibitor of metalloproteinase (*TIMP*) 2 and -4 as determined by Real Time (TaqMan) PCR in canine uterus during different times of gestation. Two comparisons are presented for each gene: First, the pre-implantation (Pre-imp.) stage was compared to the inter-placental sites during post-implantation (Post-imp.) and mid-gestation (Mid-gest.), and next to the utero-placental compartments (placentation sites) of post-implantation (Post-imp.), mid-gestation (Mid-gest.) and prepartum luteolysis. Asterisks differ at ★= $p<0.05$, ★★=
55 $p<0.01$, ★★★= $p<0.001$. Numerical data are presented as geometric means $Xg \pm$ geometric standard deviation (SD).

Figure 5:

60 Representative pictures of immunohistochemical detection of collagen 1 (COL1) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the
65 connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium;

70 After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel:

Collagen 1 was clearly detectable throughout pregnancy in all stromal compartments. Before implantation and placentation, strong signals were detected in stroma and myometrium of
75 both non-pregnant (A, B) and pregnant dogs (C, D). Following implantation and placentation, strong signals were localized in the connective tissue layer (stromal area) separating superficial from deep glands (E, H) as well as in myometrial (F, I) and placental stromal (G, J) compartments. Inlet in picture B shows a negative (isotype) control.

80 **Figure 6:**

Representative pictures of immunohistochemical detection of extracellular matrix protein 1 (ECM1) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-
85 Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J) : placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead =
90 glandular epithelium; asterisk = blood vessel;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel;

95 ECM1 was weakly detectable in luminal epithelium in the non-pregnant group (A, B). In early pregnancy, strong signals were detectable in luminal epithelium and in superficial and deep glands (C, D). During mid-gestation (E) and at prepartum luteolysis (H), ECM1 was weakly detectable in glandular epithelium of the superficial glands. At prepartum luteolysis, ECM was strongly present in the connective tissue layer separating superficial glands from
100 deep glands (H). In mid-gestation (F) and at pre-partum luteolysis (I), ECM1 was detectable

in the glandular epithelium of deep glands. Placental localization of ECM1 was in cytotrophoblast, shown at mid-gestation (G) and prepartum luteolysis (J). Inlet in picture B shows respective negative (isotype) control.

105 **Figure 7:**

Representative pictures of immunohistochemical detection of cell adhesion molecule connexin 43 (Cx43) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J) : placental compartments.

115 Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel; D = decidual cell; S = syncytiotrophoblast; C = cytotrophoblast

120 Cx43 was present in epithelial compartments and myometrial compartments. The luminal epithelium (A, C) and the epithelial cells of superficial and deep glands (A-F, H, I) stained strongly for Cx43. Additionally, the protein was present in tunica intima of blood vessels in all samples. Placental Cx43 localization was diffusely detected in cyto- and syncytiotrophoblast and decidual cells (G, J), additionally strong signals for Cx43 were detected in the intima and media of placental blood vessels. Inlet in picture B shows respective negative (isotype) control.

Figure 8:

130 Representative pictures of immunohistochemical detection of tissue inhibitor of metalloproteinases 2 (TIMP2) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); A and C:

luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel; D = decidual cell; S = syncytiotrophoblast; C = cytotrophoblast

In the non-pregnant group, TIMP2 was weakly detectable in the deep glands (B). In early pregnancy, TIMP2 was clearly detectable in luminal (C) and superficial (C) epithelium, deep glands (D) and myometrial compartments (D). At mid-gestation, TIMP2 was detectable in the glandular epithelium of the superficial glands (E), but not in the stroma of the connective tissue layer separating superficial from deep glands (E). At partum luteolysis, TIMP2 signals appeared weaker (compared to mid-gestation) in the luminal epithelium of the superficial glands; additionally, strong signals were detected in the connective tissue layer (H). At mid-gestation (F) TIMP2 appeared in the *tunica media* of blood vessels in the myometrium. At midgestation (F) and at partum luteolysis (I), TIMP2 was present in the myometrium and in deep glands. In placenta at midgestation (G) and at pre-partum luteolysis (J), TIMP2 was present in cyto- and syncytiotrophoblast, decidual cells and the intima of blood vessels. In placenta the endothelium appeared to stain strongest for TIMP2. Inlet in picture B shows the respective negative (isotype) control.

Figure 9:

Representative pictures of immunohistochemical detection of tissue inhibitor of metalloproteinases 4 (TIMP4) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at partum luteolysis (H, I, J); (A and C): luminal surface area; (B, D, F and I): deep glands area at the border of the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; asterisk = blood vessel;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel;

Before trophoblast attachment (*i.e.* at early pregnancy), and in non-pregnant controls, TIMP4 appeared weakly present in the surface epithelium (A, C). Also superficial epithelium stained weakly throughout pregnancy. Stronger signals were seen in the epithelium of deep glands, in particular in non-pregnant dogs (B) and at mid-gestation and in the prepartum luteolysis group (F, I). TIMP4 was strongly detected at prepartum luteolysis in the connective tissue layer separating superficial from deep glands (H). In the placenta, shown at mid-gestation and prepartum luteolysis, TIMP4 was present in the cytotrophoblast (G, J). Inlet in picture B shows respective negative (isotype) control.

Supplemental Files

Supplemental File 1

Immunohistochemical detection of collagen 3 (COL3) in the canine uterus and utero-placental (Ut-Pl) compartments at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; asterisk = blood vessel;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel; open arrowhead = fetal stroma. Inlet in picture B shows the respective negative (isotype) control.

Supplemental File 2:

Immunohistochemical detection of collagen 4 (COL4) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; asterisk = blood vessel;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands (*i.e.*, glands above connective layer); solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel. Inlet in picture B shows the respective negative (isotype) control.

Supplemental File 3

Immunohistochemical detection of TIMP2 and TIMP4 on consecutive slides. Four consecutive slides were prepared from placental compartments of the prepartum luteolysis group. Of these, IHC staining was performed against TIMP2 (A), TIMP4 (B), Vimentin (C) and pan-Cytokeratin (D). (A-D): open arrows = decidual cells; solid arrows = cytotrophoblast; solid arrowhead = intima of maternal blood vessels; open arrowhead = syncytiotrophoblast.

Supplemental File 4

Immunohistochemical detection of alpha smooth muscle actin (aSMA) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy; in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer; (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; asterisk = blood vessel;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel. Inlet in picture B shows the respective negative (isotype) control.

Supplemental File 5

Immunohistochemical detection of fibronectin 1 (FN1) in the canine uterus and utero-placental (Ut-Pl) compartments at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel; D = decidual cell; S = syncytiotrophoblast; C = cytotrophoblast. Inlet in picture B shows the respective negative (isotype) control.

Supplemental File 6

Immunohistochemical detection of laminin alpha 2 (LAMA2) in the canine uterus and utero-placental (Ut-Pl) (the whole thickness of the uterine wall, *i.e.*, uterus with adjacent placenta) compartments at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective layer (superficial glands, so-called glandular chambers); (G and J) : placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; asterisk = blood vessel;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel. Inlet in picture B shows the respective negative (isotype) control.

Supplemental File 7

Immunohistochemical detection of cell adhesion molecule connexin 26 (Cx26) in the canine uterus and utero-placental (Ut-Pl) compartments at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border to the myometrium; (E and H): area above the connective tissue layer; (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel; D = decidual cell; S = syncytiotrophoblast; C = cytotrophoblast. Inlet in picture B shows the respective negative (isotype) control.

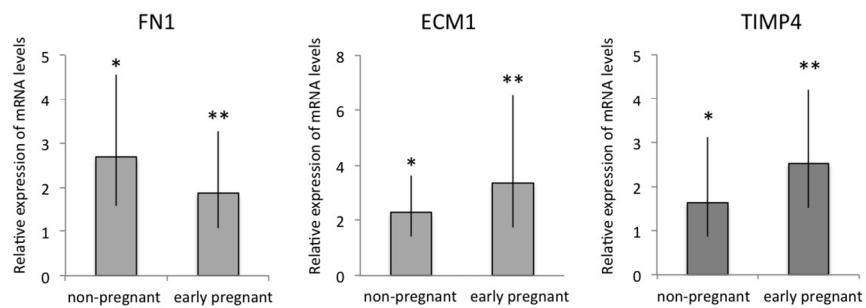


Figure 1

Expression of selected extracellular matrix (ECM) genes in the canine pre-implantation uterus (days 10-12 of pregnancy) affected by the presence of free-floating embryos, compared with gene expression in their non-pregnant counterparts as determined by Real Time (TaqMan) PCR. FN1= fibronectin 1; ECM1= extracellular matrix protein 1, TIMP4 = tissue inhibitor of metalloproteinases 4. An unpaired, two-tailed Student's t-test was applied. $P < 0.05$ was defined as significant. Bars with different asterisks differ at $P = 0.01$. Numerical data are presented as geometric means $X_g \pm$ geometric standard deviation (SD).

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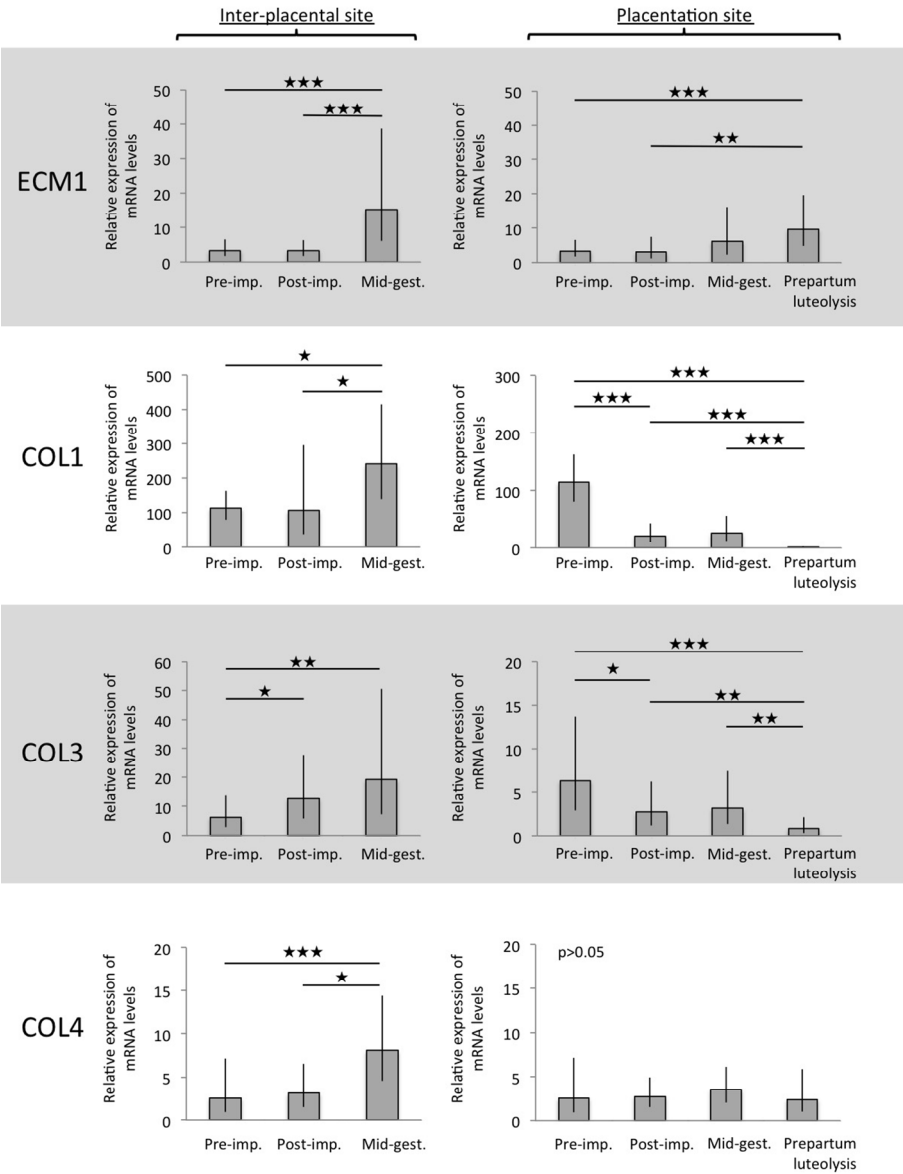
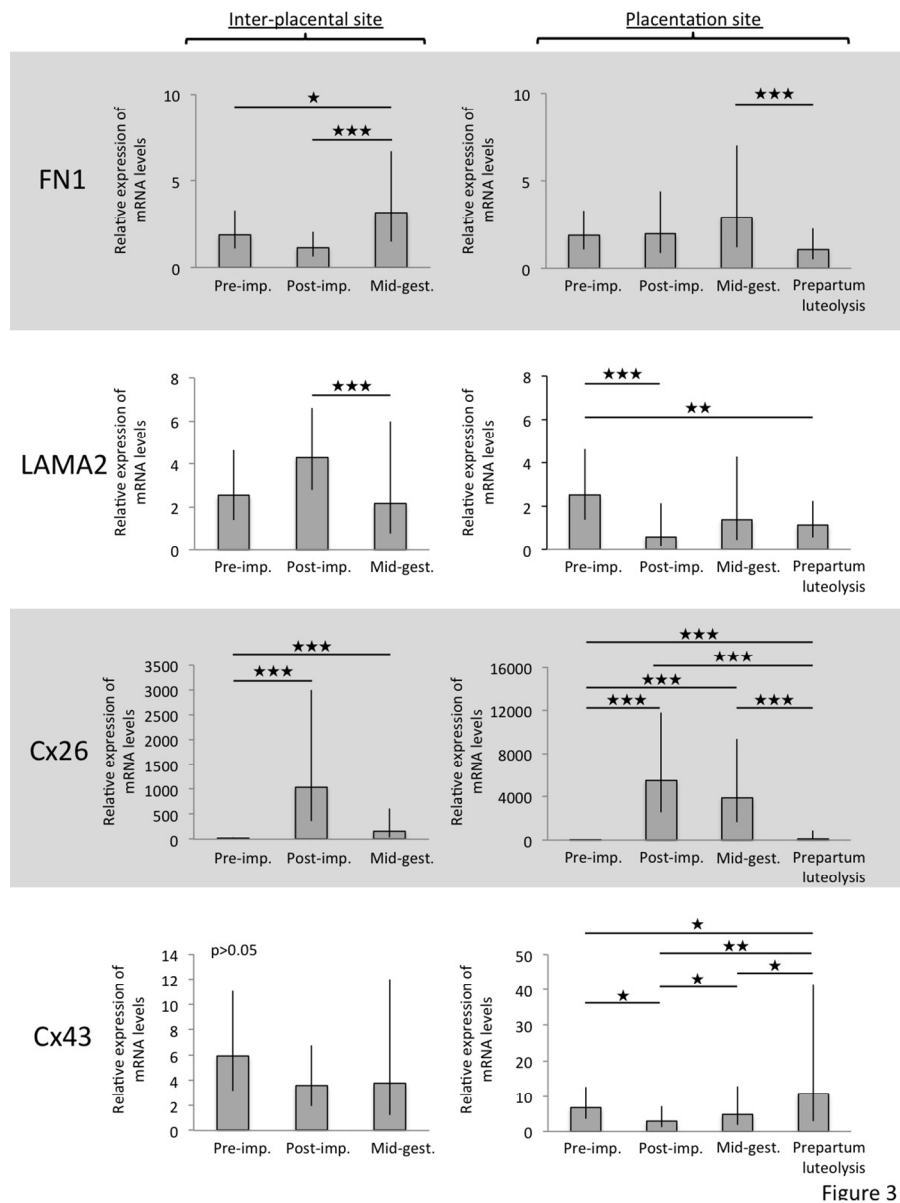


Figure 2

Expression of extracellular matrix protein 1 (ECM1), collagen (COL) 1, -3 and -4 as determined by Real Time (TaqMan) PCR in the canine uterus during different times of pregnancy. Two comparisons are presented for each gene: First, the pre-implantation (Pre-imp.) stage was compared to the inter-placental sites during post-implantation (Post-imp.) and mid-gestation (Mid-gest.), and next to the utero-placental compartments (placentation sites) of post-implantation (Post-imp.), mid-gestation (Mid-gest.) and prepartum luteolysis. Asterisks indicate ★ = $p < 0.05$, ★★ = $p < 0.01$, ★★★ = $p < 0.001$. Numerical data are presented as geometric means \pm geometric standard deviation (SD).



Expression of fibronectin 1 (FN1), laminin alpha 2 (LAMA2), connexin (Cx) 26 and -43 as determined by Real Time (TaqMan) PCR in canine uterus during different times of pregnancy. Two comparisons are presented for each gene: First, the pre-implantation (Pre-imp.) stage was compared to inter-placental sites during post-implantation (Post-imp.) and mid-gestation (Mid-gest.), and next to the utero-placental compartments (placentation sites) of post-implantation (Post-imp.), mid-gestation (Mid-gest.) and prepartum luteolysis. Asterisks differ at ★ = $p < 0.05$, ★★ = $p < 0.01$, ★★★ = $p < 0.001$. Numerical data are presented as geometric means $\bar{x} \pm$ geometric standard deviation (SD).

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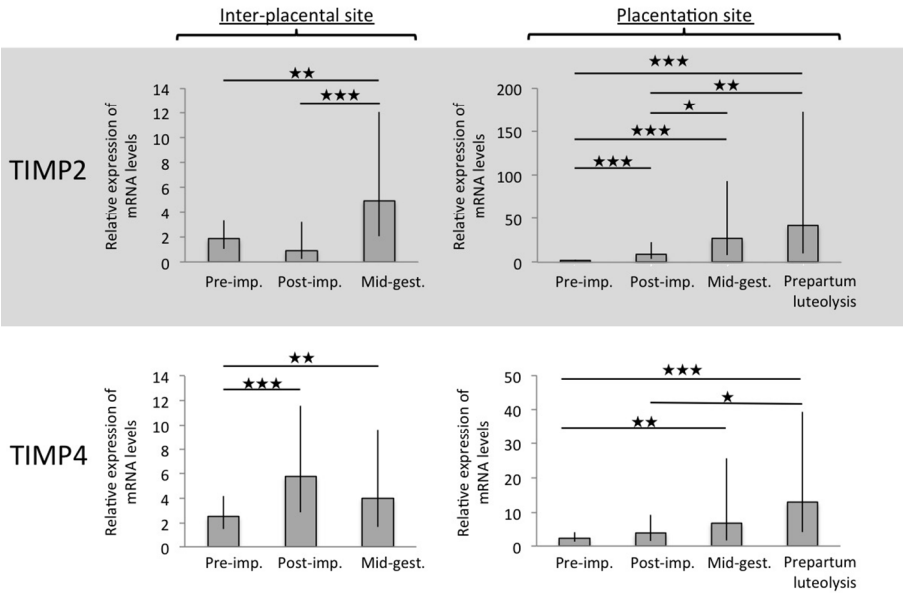


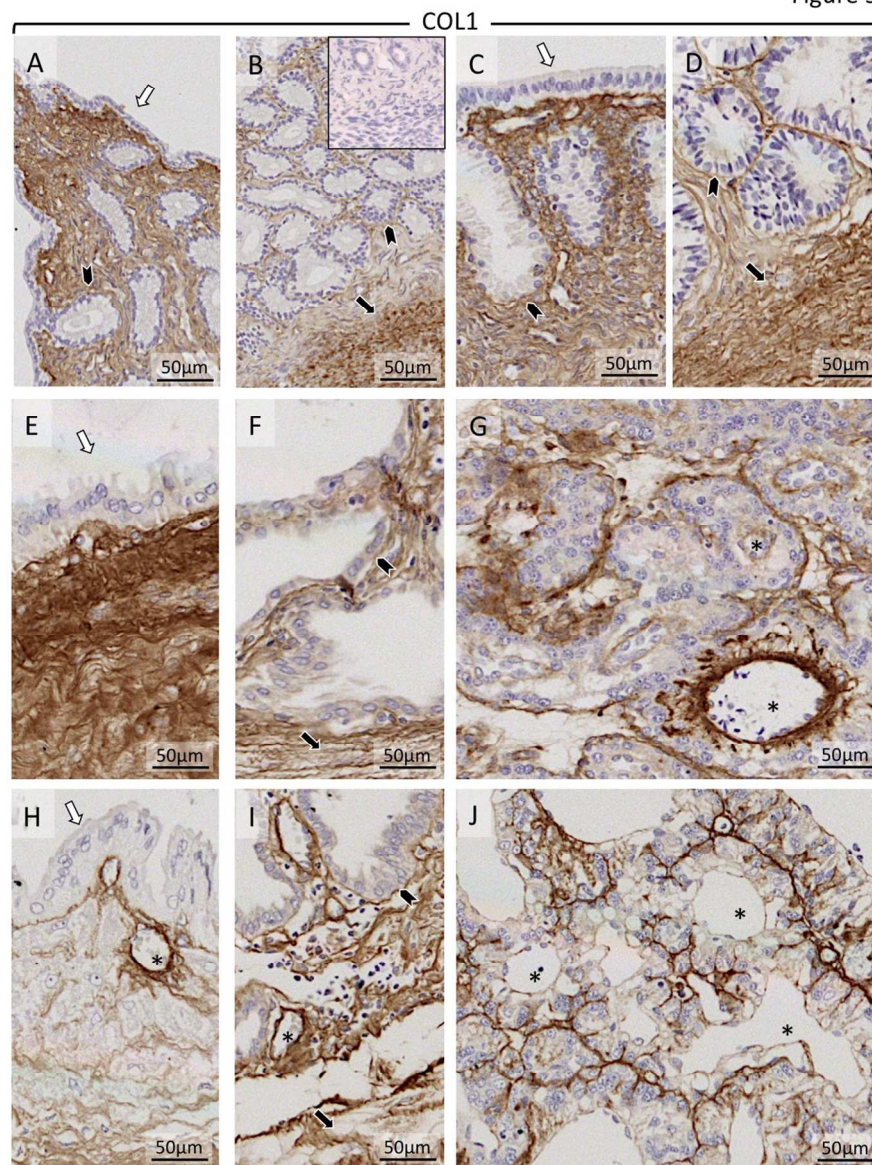
Figure 4

Expression of tissue inhibitor of metalloproteinase (TIMP) 2 and -4 as determined by Real Time (TaqMan) PCR in canine uterus during different times of gestation. Two comparisons are presented for each gene:

First, the pre-implantation (Pre-imp.) stage was compared to the inter-placental sites during post-implantation (Post-imp.) and mid-gestation (Mid-gest.), and next to the utero-placental compartments (placentation sites) of post-implantation (Post-imp.), mid-gestation (Mid-gest.) and prepartum luteolysis.

Asterisks differ at ★ = p<0.05, ★★ = p<0.01, ★★★ = p<0.001. Numerical data are presented as geometric means Xg ± geometric standard deviation (SD).

Figure 5



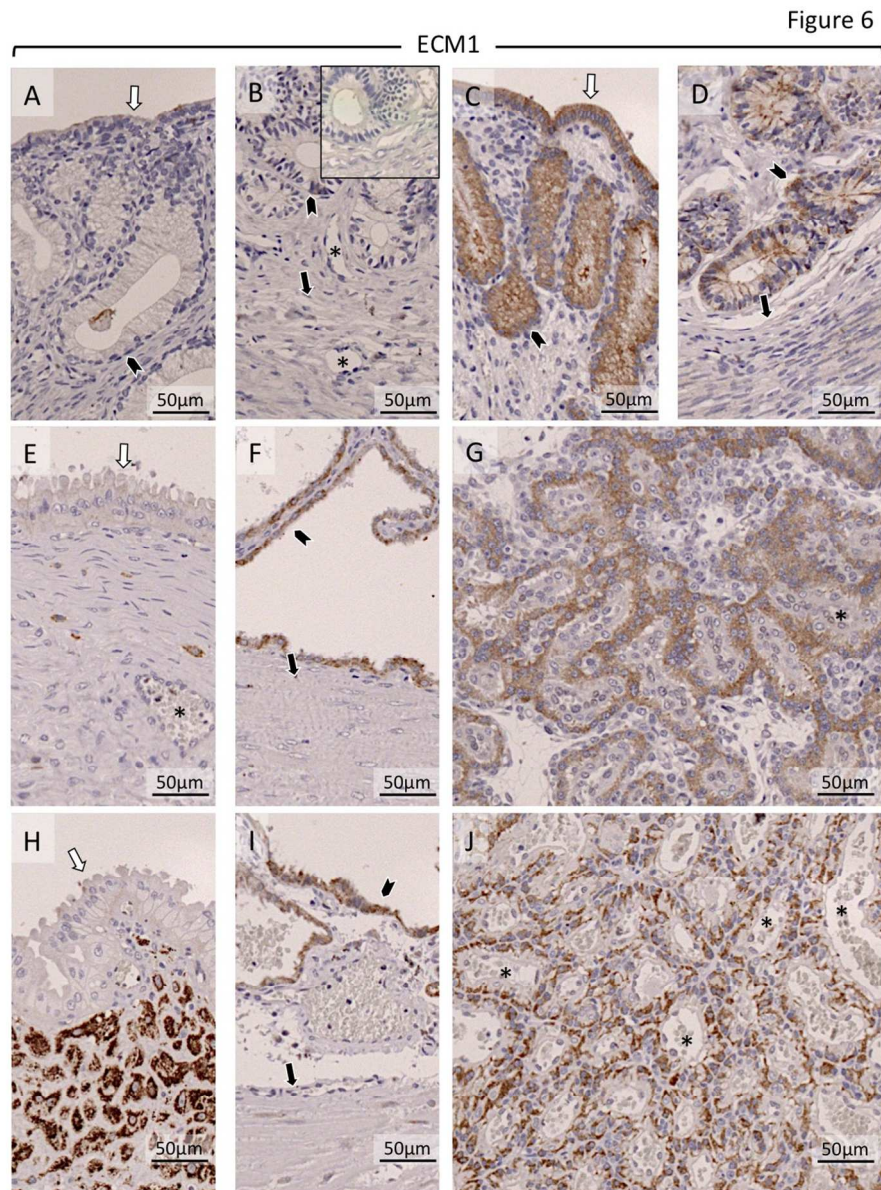
Representative pictures of immunohistochemical detection of collagen 1 (COL1) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel:

Collagen 1 was clearly detectable throughout pregnancy in all stromal compartments. Before implantation

and placentation, strong signals were detected in stroma and myometrium of both non-pregnant (A, B) and pregnant dogs (C, D). Following implantation and placentation, strong signals were localized in the connective tissue layer (stromal area) separating superficial from deep glands (E, H) as well as in myometrial (F, I) and placental stromal (G, J) compartments. Inlet in picture B shows a negative (isotype) control.

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Representative pictures of immunohistochemical detection of extracellular matrix protein 1 (ECM1) in the canine uterus and utero-placental compartments (Ut-PI) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-PI units during mid-gestation (E, F, G), and in Ut-PI compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J) : placental compartments.

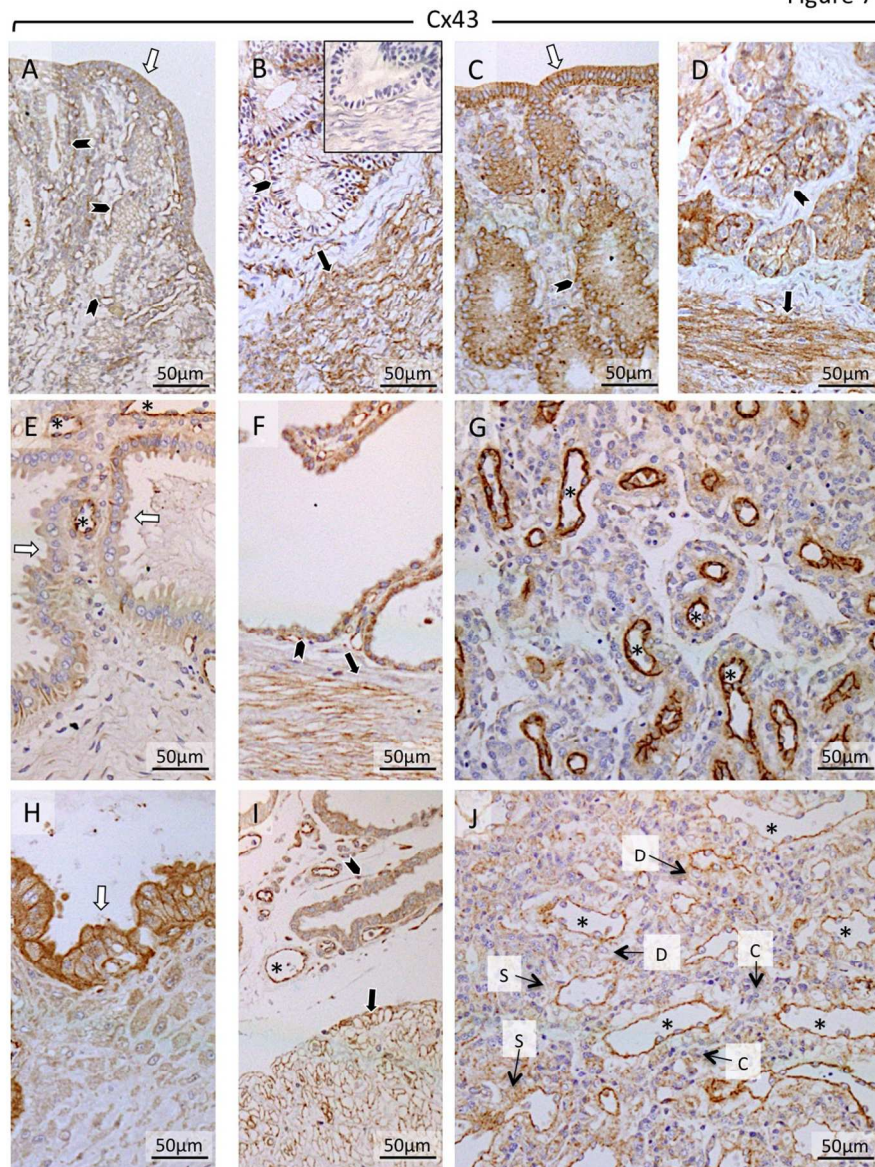
Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; asterisk = blood vessel;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel;

ECM1 was weakly detectable in luminal epithelium in the non-pregnant group (A, B). In early pregnancy, strong signals were detectable in luminal epithelium and in superficial and deep glands (C, D). During mid-gestation (E) and at prepartum luteolysis (H), ECM1 was weakly detectable in glandular epithelium of the superficial glands. At prepartum luteolysis, ECM was strongly present in the connective tissue layer separating superficial glands from deep glands (H). In mid-gestation (F) and at pre-partum luteolysis (I), ECM1 was detectable in the glandular epithelium of deep glands. Placental localization of ECM1 was in cytotrophoblast, shown at mid-gestation (G) and prepartum luteolysis (J). Inlet in picture B shows respective negative (isotype) control.

423x564mm (72 x 72 DPI)

Figure 7



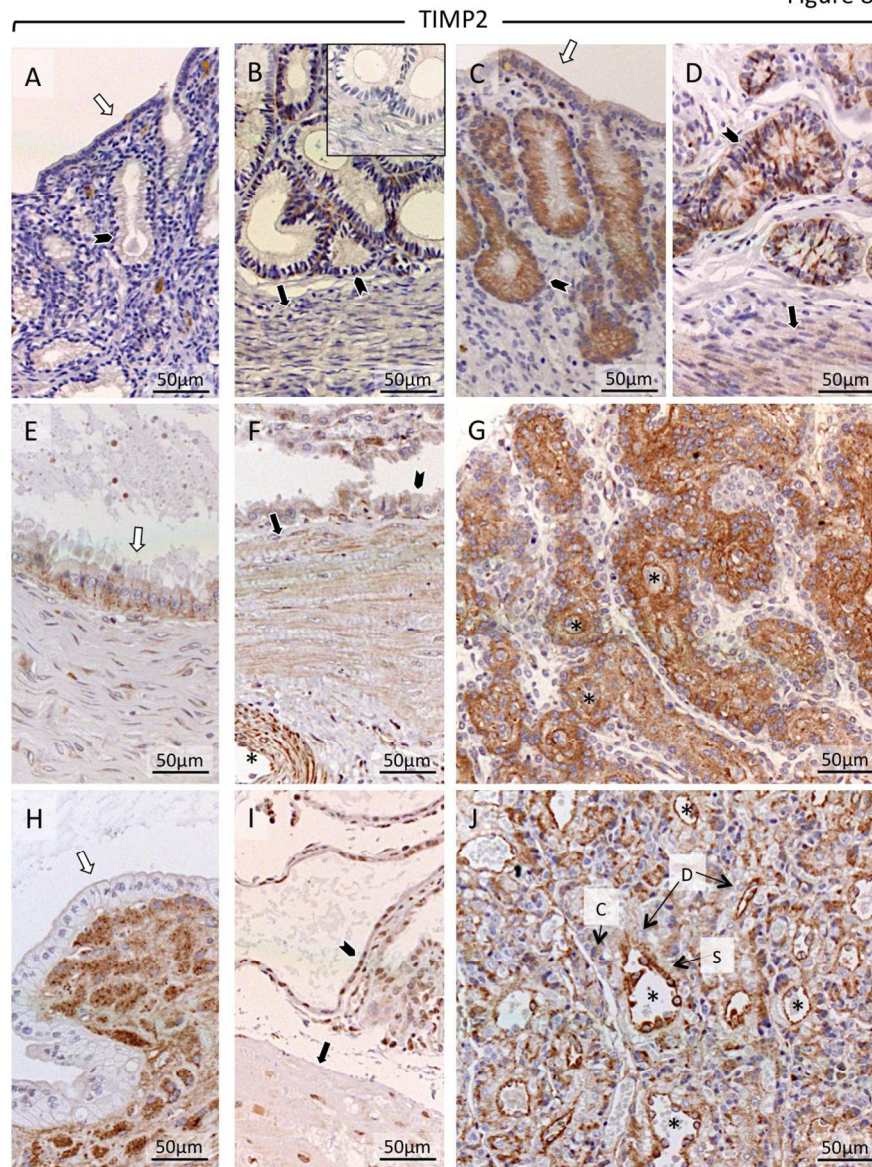
Representative pictures of immunohistochemical detection of cell adhesion molecule connexin 43 (Cx43) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J) : placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel; D = decidual cell; S = syncytiotrophoblast; C = cytotrophoblast
Cx43 was present in epithelial compartments and myometrial compartments. The luminal epithelium (A, C)

and the epithelial cells of superficial and deep glands (A-F, H, I) stained strongly for Cx43. Additionally, the protein was present in tunica intima of blood vessels in all samples. Placental Cx43 localization was diffusely detected in cyto- and syncytiotrophoblast and decidual cells (G, J), additionally strong signals for Cx43 were detected in the intima and media of placental blood vessels. Inlet in picture B shows respective negative (isotype) control.

423x564mm (72 x 72 DPI)

Figure 8



Representative pictures of immunohistochemical detection of tissue inhibitor of metalloproteinases 2 (TIMP2) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); A and C: luminal part of uterus with uterine surface area; (B, D, F and I): deep glands area at the border with the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel; D = decidual cell; S = syncytiotrophoblast; C = cytotrophoblast
In the non-pregnant group, TIMP2 was weakly detectable in the deep glands (B). In early pregnancy, TIMP2

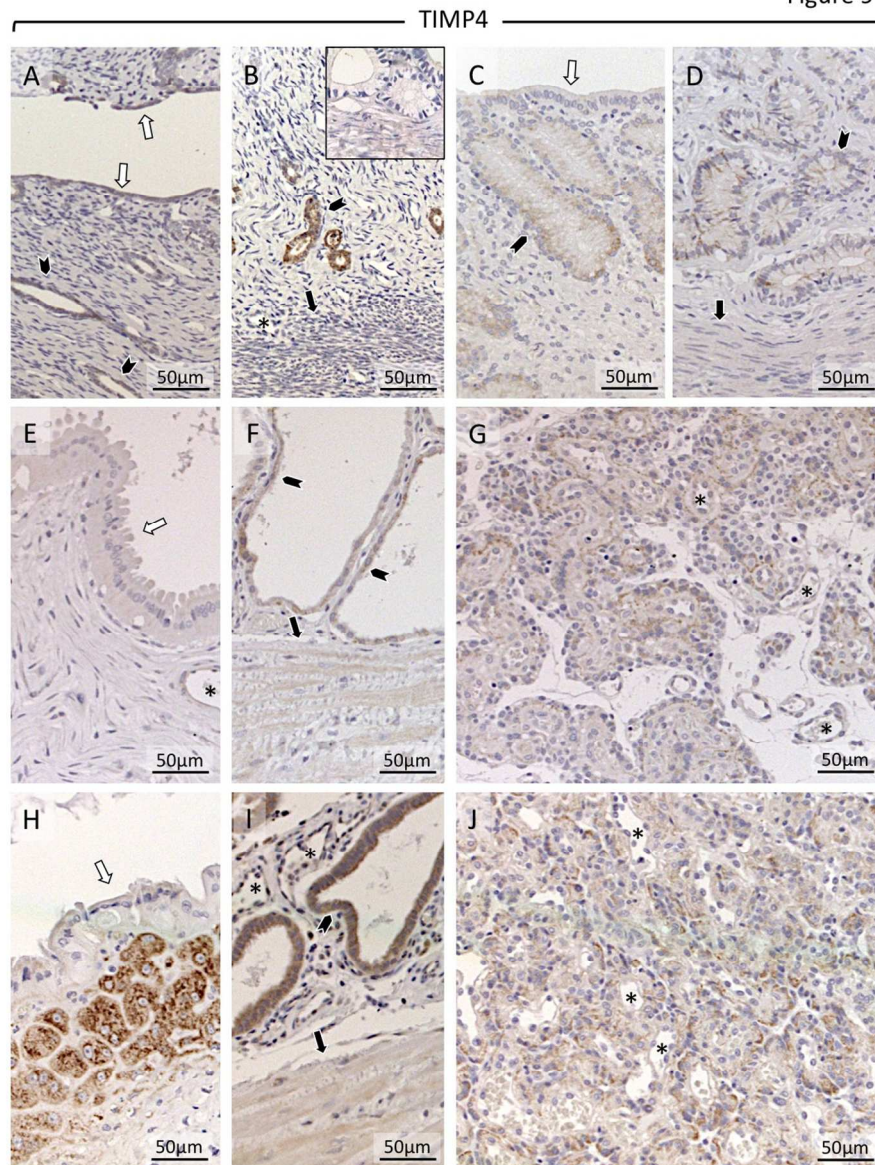
was clearly detectable in luminal (C) and superficial (C) epithelium, deep glands (D) and myometrial compartments (D). At mid-gestation, TIMP2 was detectable in the glandular epithelium of the superficial glands (E), but not in the stroma of the connective tissue layer separating superficial from deep glands (E).

At prepartum luteolysis, TIMP2 signals appeared weaker (compared to mid-gestation) in the luminal epithelium of the superficial glands; additionally, strong signals were detected in the connective tissue layer (H). At mid-gestation (F) TIMP2 appeared in the tunica media of blood vessels in the myometrium. At midgestation (F) and at prepartum luteolysis (I), TIMP2 was present in the myometrium and in deep glands.

In placenta at midgestation (G) and at pre-partum luteolysis (J), TIMP2 was present in cyto- and syncytiotrophoblast, decidual cells and the intima of blood vessels. In placenta the endothelium appeared to stain strongest for TIMP2. Inlet in picture B shows the respective negative (isotype) control.

423x564mm (72 x 72 DPI)

Figure 9



Representative pictures of immunohistochemical detection of tissue inhibitor of metalloproteinases 4 (TIMP4) in the canine uterus and utero-placental compartments (Ut-Pl) at selected time points during pregnancy: in the non-pregnant canine uterus at early diestrus (A, B), early-pregnant (pre-implantation) canine uterus (C, D), Ut-Pl units during mid-gestation (E, F, G), and in Ut-Pl compartments at prepartum luteolysis (H, I, J); (A and C): luminal surface area; (B, D, F and I): deep glands area at the border of the myometrium; (E and H): area above the connective tissue layer (superficial glands, so-called glandular chambers); (G and J): placental compartments.

Non-pregnant (A-B), and before implantation and placentation (C-D): open arrows = surface (luminal) epithelium; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium; asterisk = blood vessel;

After implantation and placentation (E-J): open arrows = glandular epithelium of superficial glands; solid arrows = myometrium (circular layer); solid arrowhead = glandular epithelium of deep glands; asterisk = blood vessel;

Before trophoblast attachment (i.e. at early pregnancy), and in non-pregnant controls, TIMP4 appeared weakly present in the surface epithelium (A, C). Also superficial epithelium stained weakly throughout pregnancy. Stronger signals were seen in the epithelium of deep glands, in particular in non-pregnant dogs (B) and at mid-gestation and in the prepartum luteolysis group (F, I). TIMP4 was strongly detected at prepartum luteolysis in the connective tissue layer separating superficial from deep glands (H). In the placenta, shown at mid-gestation and prepartum luteolysis, TIMP4 was present in the cytotrophoblast (G, J). Inlet in picture B shows respective negative (isotype) control.

423x564mm (72 x 72 DPI)

Primer	Accession number	Primer Sequence		Product length (bp)
extracellular matrix protein 1 (<i>ECM1</i>)	XM_845921.4	Forward	5'-CAG TCT GGC TTC TCC CAC CTT A-3'	99
		Reverse	5'-GCG GTT TGT GTG GCT GTG A-3'	
		TaqMan Probe	5'-AGA CTA GAT ATT CCC GCT GCT GCC GCT-3'	
connexin 26 (<i>Cx26</i>)	AJ439693.1	Forward	5'-CCA CTA CTT CCC CAT CTC TCA CA -3'	98
		Reverse	5'-TCC GGT AGG CGA CAT GCA T -3'	
		TaqMan Probe	5'-CCG ACT CTG GGC TCT GCA GCT GAT C-3'	
connexin 43 (<i>Cx43</i>)	AY462223	Forward	5'-AAA AGA GAA CCC TGC CCT CAT C-3'	91
		Reverse	5'-AGG ACA CGA CCA GCA TGA AGA-3'	
		TaqMan Probe	5'-ACT GCT TCC TCT CTC GCC CCA CG-3'	
fibronectin 1 (<i>FN1</i>)	XM_014110981	Forward	5'-CAC GCC GAA CTA CGA TGC-3'	95
		Reverse	5'-TGC GAT ACA TGA CCC CTT-3'	
		TaqMan Probe	5'-AAG TTT GGA TTT TGC CCC ATG GCC-3'	
laminin alpha 2 (<i>LAMA2</i>)	XM_014113700.1	Forward	5'-AAA CCG GCT CAC GAT TGA G-3'	99
		Reverse	5'-AGT TGA ACG GTG GCG AAG T-3'	
		TaqMan Probe	5'-CCT GCT CTT CTA CAT GGC TCG GAT CAA-3'	
glyceraldehyd 3-phosphate dehydrogenase (<i>GAPDH</i>)	AB028142	Forward:	5'-GCT GCC AAA TAT GAC GAC ATC A-3'	75
		Reverse:	5'-GTA GCC CAG GAT GCC TTT GAG-3'	
		TaqMan Probe	5'-TCC CTC CGA TGC CTG CTT CAC TAC CTT-3'	
tissue inhibitor of matrix metalloproteinase-2 (<i>TIMP2</i>)	AF188489	Forward	5'-CAT AGG TAC CAG ATG GGC TGT GA-3'	95
		Reverse	5'-CAG TCC ATC CAG AGG CAC TCA-3'	
		TaqMan Probe	5'-TGA TCC CGT GCT ATA TCT CGT CTC CGG-3'	
tissue inhibitor of matrix metalloproteinase-2 (<i>TIMP4</i>)	NM_001314106	Forward	5'-CTG TGG CTG CCA AAT TAC CA-3'	103
		Reverse	5'-CCC ATA GAG CTT CCG TTC CA-3'	
		TaqMan Probe	5'-ACC ATC TCA GCC CCT AAC GAG TGC CTC-3'	

Table 1

Antibody	Company	Reference Number	Antigen-Retrieval	Dilution
collagen I (COL1)	Rockland	31680	PIER 60min	1:800
collagen III (COL3)	Quartett	292302	PIER 45min	1:200
collagen IV (COL4)	Quartett	250424	PIER 60min	1:500
laminin 2 alpha (LAMA2)	Bioss Antibodies	bs-8561R	PIER 90min	1:300
fibronectin 1 (FN1)	Novus Biologicals	NBP1-91258	PIER 30min	1:300
alpha-smooth-muscle-actin (aSMA)	Dako	M0851	HIER	1:100
connexin 26 (Cx26)	Thermo-fisher	QC215178	PIER 5min	1:200
connexin 43 (Cx43)	Abcam	AB11370	PIER 15min	1:750
tissue inhibitor of matrix metalloproteinase-2 (TIMP2)	Merck Millipore	AB2965	HIER	1:1500
tissue inhibitor of matrix metalloproteinase-2 (TIMP4)	LifeSpan Biosciences	LS-C116809	HIER	1:750
anti-vimentin clone vim3B4 (VIM)	Dako	M7020	HIER	1:100
anti-cytokeratin (pan-CYT)	Dako	Z0622	HIER	1:700
extracellular matrix protein 1 (ECM1)	Proteintech	11521-1-AP	HIER	1:200
biotinylated goat anti-rabbit IgG (H+L)	Vector Laboratories	BA-1000	-	1:100
biotinylated horse anti-mouse IgG (H+L)	Vector Laboratories	BA-2000	-	1:100

Table 2

	COL1	COL3	COL4	FN1	ECM1	LAMA2	Cx26	Cx43	TIMP2	TIMP4	aSMA
NON-PREGNANT											
ENDOMETRIUM											
Luminal epithelium	-	-	-	+	+	+	+/-	+	+/-	+/-	-
Stroma	+	+	-	+	-	+	-	-	-	-	+
Glandular epithelium	-	-	-	+	-	+	+/-	+	+/-	+	-
Blood vessels	-	-	+	-	-	+	-	+	+	-	++
MYOMETRIUM											
Smooth muscle cells	-	-	-	-	-	-	+	+	+/-	-	++
Stroma	+	+	+	+	-	+	-	-	+	-	-
Blood vessels	-	-	+	-	-	+	-	+	+	-	++
PRE-ATTACHMENT											
ENDOMETRIUM											
Luminal epithelium	-	-	-	+	++	+	+	+	++	+/-	-
Stroma	+	+	-	+	-	+	-	-	-	-	-
Glandular epithelium	-	-	-	+	+	+	+	+	++	+	-
Blood vessels	-	-	+	-	-	+	-	+	+	-	+
MYOMETRIUM											
Smooth muscle cells	-	-	-	-	-	+	+	+	+	-	+
Stroma	+	+	+/-	+	-	+	-	-	+	-	-
Blood vessels	-	-	+	-	-	+	-	+	+	-	++
MID-GESTATION											
PLACENTA											
Cytotrophoblast cells	-	-	-	+	+	-	+	+/-	+	+	-
Syncytiotrophoblast	-	-	-	+	-	-	+	+/-	+	-	-
Decidual cells	-	-	-	+	-	-	+	+/-	+	-	++
Blood vessels	-	-	+	+	-	+	+	++	+	-	+
Stroma	+	+	-	+	-	+	-	-	-	-	-
ENDOMETRIUM											
Epithelium of superficial glands	-	-	-	+	+	+	++	+	+	+/-	-
Connective tissue layer separating superficial from deep glands	+	+	-	+	-	++	-	+	+/-	-	-
Epithelium of deep glands	-	-	-	+	+	+	+	+	+	+	-
Blood vessels	-	-	+	+	-	+	+/-	++	+	-	+
MYOMETRIUM											
Smooth muscle cells	-	-	-	-	-	+	+	+	+	-	+
Blood vessels	-	-	+	-	-	+	-	+	+	-	++
Stroma	+	+	+	+	-	+	-	-	+	+	-
PREPARTUM LUTEOLYSIS											
PLACENTA											
Cytotrophoblast cells	-	-	-	+	+	-	++	+	+	+	-
Syncytiotrophoblast	-	-	-	+	-	-	+	+	+	-	-
Decidual cells	-	-	-	+	-	-	+	+	+	-	++
Blood vessels	-	-	+	+	-	+	+/-	++	++	-	++
Stroma	+	+	-	+	-	+	-	-	-	-	-
ENDOMETRIUM											
Epithelium of superficial glands	-	-	-	+	+	+	++	+	+/-	+/-	-
Connective tissue layer separating superficial from deep glands	+	+	-	+	++	+	+	+	++	++	-

Epithelium of deep glands	-	-	-	+	+	+	+	+	+	++	-
Blood vessels	-	-	+	-	-	+	+	++	+	-	++
MYOMETRIUM											
Smooth muscel cells	-	-	-	-	-	+	+	+	+	-	+
Blood vessels	-	-	+	-	-	+	+	+	+	-	+
Stroma	+	+	+/-	+	-	+	-	-	+	+	-

Table3